

Preliminary Amendment

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- 2) The chimeric expression promoter according to claim 1, wherein said first plant promoter originates from the Commelina Yellow Mottle Virus and said second plant promoter originates from the Cassava Vein Mosaic Virus.
  - 3) The chimeric promoter of claim 1 wherein said nucleic acid sequences originate from the intergenic regions of said first and second promoters.
  - 4) The chimeric promoter of claim 1, wherein it comprises at least a part of a nucleic acid sequence with SEQ ID No. 1 fused to at least a part of a nucleic acid sequence with SEQ ID No. 2.
  - 5) The chimeric promoter of claim 1, wherein the nucleic acid sequence of said chimeric promoter consists of a sequence selected from the group consisting of the sequences with SEQ ID Nos. 3-7 and 19-25.
  - 6) A chimeric expression promoter comprising a promoter of viral origin, of which a part consists of an exogenous element which promotes expression in plant green tissues.
  - 7) The chimeric expression promoter of claim 6, wherein said exogenous promoter element is also of viral origin.
  - 8) The chimeric expression promoter of claim 7, wherein said promoter of viral origin originates from the Commelina Yellow Mottle Virus.
  - 9) The chimeric expression promoter of claim 7, wherein said exogenous promoter element originates from the Cassava Vein Mosaic Virus.
  - 10) The chimeric expression promoter of claim 6, wherein the exogenous element replaces an endogenous vascular tissue expression promoter of viral origin.

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- 11) The chimeric promoter of claim 1 or 6, wherein it further comprises at least one “endosperm like” box.
  - 12) The chimeric promoter of claim 1 or 6, wherein it further comprises at least one “as1 like” box operably linked to a plant green tissue expression GT promoter element.
  - 13) The chimeric promoter of claim 1 or 6, wherein it further comprises at least one “as1” box operably linked to a green tissue expression GT promoter element.
  - 14) The chimeric promoter of claim 1 or 6, wherein it further comprises at least one “as2” box operably linked to a plant green tissue expression GT promoter element.
  - 15) The chimeric promoter of claim 1 or 6, wherein said “as1 like”, “as1”, and “as2” boxes are operably linked upstream or downstream of the plant green tissue expression GT promoter element.
  - 16) The chimeric promoter of claim 1 or 6, wherein said “as1 like”, “as1”, and “as2” boxes are operably linked in 5’>3’ or 3’>5’ orientation.
  - 17) The chimeric promoter of claim 1 or 6, wherein it comprises at least one “as2/as2/as2” box in 5’>3’ or 3’>5’ orientation.
  - 18) The chimeric promoter of claim 6, wherein it comprises at least one sequence selected from the group consisting of sequences with SEQ ID Nos. 3-7 and 19-25.
  - 19) An expression cassette comprising at least one nucleic acid sequence derived from a first plant promoter comprising a plant vascular expression promoter region, said plant vascular expression promoter region being replaced with a nucleic acid sequence derived from a second plant promoter and comprising a plant green tissue expression promoter region, said at least one

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sequence being operably linked to a nucleic acid sequence or gene coding for a polypeptide to be produced, said nucleic acid sequence or gene itself operably linked to a transcription termination nucleic acid sequence.

20) The expression cassette of claim 19, wherein said first plant promoter originates from the Commelina Yellow Mottle Virus and said second plant promoter originates from the Cassava Vein Mosaic Virus.

21) The expression cassette of claim 19, wherein it comprises at least a part of a nucleic acid sequence with SEQ ID No. 1 fused to at least a part of a nucleic acid sequence with SEQ ID No. 2.

22) The expression cassette of claim 19, wherein said first or second promoter comprises a sequence selected from the group consisting of the sequences with SEQ ID Nos. 3-7 and 19-25.

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32) A guide deoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to claim 1, 6, or 49 wherein said sequence is selected from the group consisting of SEQ ID Nos. 15-18.

33) Guide deoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ ID No. 16.

36) A vector comprising a promoter, or a promoter nucleic acid sequence, which initiates transcription of a nucleic acid sequence[, or gene,] coding for a polypeptide, wherein said promoter or said isolated promoter nucleic acid sequence comprises a chimeric expression promoter or a promoter nucleic acid sequence according to claim 1, 6, or 49.

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37) The vector of claim 36, wherein said vector is selected from the group consisting of the binary vectors pMRT1152, pMRT1171, pMRT1172, pMRT1185, pMRT1186, pMRT1187, pMRT1188, pMRT1182, pMRT1245, pMRT1246, pMRT1247, pMRT1248, pMRT1249, pMRT1250, pMRT1251, pMRT1252, pMRT1253 and pMRT1254.

38) A transgenic plant comprising stably integrated into its genome at least one promoter or at least one promoter nucleic acid sequence according to claim 1, 6, or 49.

39) The transgenic plant of claim 38, wherein said plant is one selected from dicotyledonous species comprising potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, beetroot, and sunflower, or from monocotyledonous species comprising wheat, barley, oat, rice, and corn.

40) A propagule of a transgenic plant according to claim 38 or 39.

41) The transgenic plant propagule of claim 40, wherein it is a seed.

42) A cell containing a promoter or a promoter nucleic acid sequence of claim 1, 6, or 49, wherein said cell is selected from the group consisting of a plant cell, human cell, animal cell, insect cell, bacterial cell, yeast cell, fungal cell, algal cell, and microalgal cell.

43) The cell of claim 42, wherein it is a plant cell.

44) A method for expressing a nucleic acid sequence coding for a polypeptide by a cell, wherein said method comprises:

- transforming said cell with a vector comprising at least one promoter or at least one promoter nucleic acid sequence of claim 1, 6, or 49;

- culturing said cell and expressing said polypeptide encoded by said sequence in said

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cell.

45) The method of claim 44, wherein said cell is a prokaryotic or an eukaryotic cell.

46) The method of claim 44 or 45, wherein said cell is a cell selected from the group consisting of bacterial cells, fungal cells, yeast cells, insect cells, human cells, animal cells, algal cells, microalgal cells and plant cells.

47) The method of claim 46, wherein said cell is a plant cell.

48) A method for manufacturing a transgenic plant of claim 38, or a propagule of claim 40, wherein said method comprises[ the steps consisting of]:

- transforming a plant cell with a vector comprising at least one promoter or at least one promoter nucleic acid sequence of claim 1, 6, or 49;

- selecting said plant cell comprising integrated said promoter or said promoter nucleic acid sequence;

- propagating said selected plant cell by culture or by regeneration of whole chimeric or transgenic plants.

49) An isolated promoter nucleic acid sequence, comprising a fusion of a first sequence and a second sequence, wherein said first sequence and said second sequence comprise at least a part of each of the sequences with SEQ ID No. 1 and SE ID No. 2, respectively.

50) The isolated promoter nucleic acid sequence of claim 23, wherein said first or second sequence comprises a sequence selected from the group consisting of sequences with SEQ ID Nos. 3-7 and 19-25.

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51) A directional deoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to claim 1, 6, or 23, wherein said sequence is selected from the group consisting of SEQ ID Nos. 8-14.

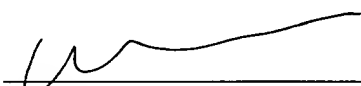
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**REMARKS**

Upon entry of this amendment, claims 1-22, 32, and 36-51 are pending. No new matter is introduced by this amendment and support for this amendment may be found in the claims as originally filed. Applicant submits that all claims are allowable as written and respectfully requests early favorable action by the Examiner.

Respectfully submitted,

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Kathleen M. Williams  
PALMER AND DODGE, LLP  
111 Huntington Avenue  
Boston, MA 02199  
(617) 239-0667